

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/658,659B TIME: 09:10:56

DATE: 11/07/2002

Input Set : A:\11926-015001.txt

Output Set: N:\CRF4\11072002\I658659B.raw

4 <110> APPLICANT: Stanton, Jr., Vincent P.

7 <120> TITLE OF INVENTION: GENE SEQUENCE VARIANCE IN GENES RELATED & C

TO FOLATE METABOLISM HAVING UTILITY IN DETERMINING THE

TREATMENT OF DISEASE 9

11 <130> FILE REFERENCE: 11926-015001

13 <140> CURRENT APPLICATION NUMBER: 09/658,659B

14 <141> CURRENT FILING DATE: 2000-09-08

16 <150> PRIOR APPLICATION NUMBER: 09/596,033

17 <151> PRIOR FILING DATE: 2000-06-15

19 <150> PRIOR APPLICATION NUMBER: 09/357,743

20 <151> PRIOR FILING DATE: 1999-07-20

22 <150> PRIOR APPLICATION NUMBER: 09/357,024

23 <151> PRIOR FILING DATE: 1999-07-19

25 <150> PRIOR APPLICATION NUMBER: 60/093,484

26 <151> PRIOR FILING DATE: 1998-07-20

28 <160> NUMBER OF SEQ ID NOS: 16

30 <170> SOFTWARE: FastSEQ for Windows Version 4.0

Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

32 <210> SEQ ID NO: 1

33 <211> LENGTH: 7224

34 <212> TYPE: DNA

35 <213> ORGANISM: Homo sapiens

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38 <221> NAME/KEY: misc_feature

39 <222> LOCATION: 194, 32/09

40 <223> OTHER INFORMATION: n = c or g

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43 <222> LOCATION: 11,36, 13,34, 3150, 5,551, 5,934

44 <223> OTHER INFORMATION: n = a or g

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47 <222> LOCATION: 284, 1252, 1699, 5\$73, 5659, 5678, 5874

48 <223> OTHER INFORMATION: n = c or t

W--> 50 <221> misc_feature

51 <222> LOCATION: 3207

52 < 223 > OTHER INFORMATION: n = g or t

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56 < 223 > OTHER INFORMATION: n = c or a

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PATENT APPLICATION: US/09/658,659B

DATE: 11/07/2002 TIME: 09:10:56

Input Set : A:\11926-015001.txt
Output Set: N:\CRF4\11072002\1658659B.raw

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							ggcaggctcg	3.0.0
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					caacatgtca			420
					ggatgagatc			480
					catgatccag			540
					tgccaggccg			600
					ccaaatccat			660
							ctgactatgg	720
							gaaaagctgc	780
					gaggtttgtg			840
					ggaaaggccg			900
							gcggggttga	960
							tgtttgcact	1020
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Total I also							ggacgatcgt tcagcntgtc	1140
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M>							chaatgcagg	
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W>							caacaccaga	1380
					aaattgtaag			1440
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							ttgctaaact	1560
					gtgtgttgcc			1620
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•							aacttetgea	2280
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							accttatecc	
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							, acataggcaa	2760
							: taggagtcat	2820
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	107	gtcaggacto	atcactcctt	ccctggatga	. aatgatttt	gttgccaagg	aaatggagag	2940
	108	attagctata	ı aggattccat	: tgttgattgg	aggagcaacc	: acttcaaaaa	cccacacage	3000

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1	W>	111	catggaagaa	tatgaagata.	ttagacaggn	ccattatgag	tctctcaagg	agaggagata	3180
,	W>	112	cttaccctta	agtcaagcca	gaaaaantng	tttccaaatg	gattggctgt	ctgaacetea	3240
		113	cccagtgaag	cccacgttta	ttgggaccca	ggtctttgaa	gactatgacc	tgcagaagct	3300
		114	ggtggactac	attgactgga	agcctttctt	tgatgtctgg	cagctccggg	gcaagtaccc	3360
		115	gaatcgaggc	tttcccaaga	tatttaacga	caaaacagta	ggtggagagg	ccaggaaggt	3420
		116	ctacgatgat	gcccacaata	tgctgaacac	actgattagt	caaaagaaac	tccgggcccg	3480
		117	gggtgtggtt	gggttctggc	cagcacagag	tatccaagac	gacattcacc	tgtacgcgga	3540
		118	ggctgctgtg	ccccaggctg	cagageceat	agccaccttc	tatgggttaa	ggcaacaggc	3600
		119	tgagaaggac	tctgccagca	cggagccata	ctactgcctc	tcagacttca	tegeteeett	3.660
		120	gcattctggc	atccgtgact	acctgggcct	gtttgccgtt	gcctgctttg	gggtagaaga	3720
		121	gctgagcaag	gcctatgagg	atgatggtga	cgactacagc	agcatcatgg	tcaaggcgct	3780
		122	gggggaccgg	ctggcagagg	cctttgcaga	agagetecat	gaaagagttc	gccgagaact	3840
		123	gtgggcctac	tgtggcagtg	agcagctgga	cgtcgcagac	ctgcgcaggc	tgcggtacaa	3900
		124	gggcatccgc	ccggctcctg	gctaccccag	ccagcccgac	cacaccgaga	agctcaccat	3960
		125	gtggagactt	gcagacatcg	agcagtctac	aggcattagg	ttaacagaat	cattagcaat	4020
		126	ggcacctgct	tcagcagtct	caggceteta	cttctccaat	ttgaagtcca	aatattttgc	4080
		127	tgtggggaag	atttccaagg	atcaggttga	ggattatgca	ttgaggaaga	acatatctgt	4140
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		129	ttttttgcct	tttttattct	tgatgatect	caaggaaata	caacctaggg	tgccttaaaa	4260
		130	ataacaacaa	caaaaaacct	gtgtgcatct	ggctgacact	tacctgatta	tggttttcga	4320
		131	agactattta	gtggaacctt	gtagaggagc	agggtcttcc	tgcagtgcct	ggaaaacagg	4380
		132	cgctgttttt	ttgggacctt	gcgtgaagag	cagtgagcag	ggttcctgtg	gtttccctgg	4440
		133	tecetetgag	atggggacag	actgaagaca	gaggtcgttt	gatttcaaag	caagtcaacc	4500
		134	tgcttttttc	tgtttttaca	gtggaatcta	ggaggccact	tagtcgtctt	tttttcctct	4560
		135	tagaagaaaa	gcctgaaact	gagttgaata	gagaagtgtg	accctgtgac	aaaatgatac	4620
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		137	gtagctctgt	tctcttcgga	agacctcgtt	ttctaaaggc	tggactaaat	ggctgcagaa	4740
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		141	taacattttt	tttgttttgt	tttgttttgg	ttttttttt	tttgagacag	agtctggctc	4980
		142	tgtcgcccag	gctggagtgc	aggggcgcaa	tctcggctca	tagcaagctc	cgcctcctgg	5040
		143	gttcatgcca	ttctcctgcc	tcagcctcca	gagtagctgg	gactacaggt	gcccgccacc	5100
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		147	ccaggattga	gcaggttctc	agctgggctc	tgatatcccg	tgcggagttg	gacaagtggg	5340
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V	v>	149	atgaagaatg	gtgccaccac	tcaagcaaca	agcctcaaac	tcanccatgt	catcttttc	5460
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V	V>	151	gatggtggtg	gcaatagtca	ggagaaggta	ncattggagt	cctggtttga	ttngaaggat	5580
							tttgttcatg		5640
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		154	aatggaaatt	ccaacagatt	ttattgaatg	aaacaagcag	gtgtttatat	ggagtagcaa	5760
		155	aggacttaaa	attaccaaat	gcttctaaat	atgaaggaga	ggttggggac	acgçacccta	5820
V	√>	156	tgtgatacca	agttttattg	tcaagacagt	gtcatggtgc	agaggtaggc	attntgagca	5880
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Input Set : A:\11926-015001.txt

		Output Set: N:\CRF4\11072002\1658659B.raw	
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		ttccaaatat gttaagggct ttaatacaaa aagcaaaaat tgtcagtgtt tggatgaaaa	6120
		aaqccttagg gcaggaaaga atctcttgag acataaagta gtaatcataa aggacaagat	6180
	162	ggttaagtca attctgttaa aactcaaggc ttatattaag caaacacttg aagtgagaag	6240
	163	atgatccaca acttgagaag acatttataa tacaaataac tgatgaagga ttcataatca	6300
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	165	atagggettt taaataaata gatgttetgt ageattggte agggaaatat gaattaggae	6420
	166	cacaatgaga ttccatttta tatccataag atttgcaaag gttgggtctg acagtaccag	6480
	167	ttgttagatc tgtagggact tgtacaacat tgtggatgtg taaacaggca ccactgcttt	6540
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	169	toccacctgt atgtccagca aactcttgca tgtggccact aggaggaatg tgtaagaatg	6660
	170	ttcatagtta catatttata atagttaata actggaaaaa gtgaaatgta tgtctgtcta	6720
		caggaaaata ggtgaataat tagatatatg tattcattct acgggatatt attcagtagt	6780
	172	ggaaatgagt gaactacagc tatacctcac aataagaatg aatctcagaa aatattaagg	6840
		aaaaaagcaa gtttgaagag accacatggg gcgtactatt tttattgagc ccaaaaacaa	6900
		gcaaaaccaa agaatatgta gtctaagcat acgtatacaa taaaactatg ctattaaaaa	6960
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		agggacagga gcacataggt agatgccaag ttatgcaget gttctggttc ctcctggtag	70/80%
		gcttacaagt/gtttactata_tgctattaat/acattatact/ttataactaa/tagataacag/	7140 persent hard
		tttttttacat attaaatatg ttctacttaa atatattata aaaaataaag gcaaagtgga	7200 return
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	200	gcatccctgg	cttgcaagga	gtagacactc	gggagctgac	caagaagttg	cgggaacagg	420
	201	ggtctctgct	ggggaagctg	gtccagaatg	gaacagaacc	ttcatccctg	ccattcttgg	480
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	205	aagagtatga	gggtctcttc	ttaagtaatg	ggcctggtga	ccctgcctcc	tatcccagtg	720
	206	togtatocac	actgagccgt	gttttatctg	agcctaatcc	ccgacctgtc	tttgggatct	780
	.207	gcctgggaca	ccagctattg	gccttagcca	ttggggccaa	gacttacaag	atgagatatg	840
	208	ggaaccgagg	ccataaccag	ccctacttat	taataaacte	tagacactac	tttctgacat	900

PATENT APPLICATION: US/09/658,659B

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	217	ttcttcccat	aacacctcat	tatgtaaccc	aggtgatacg	taatgaacgc	cccgatggtg	1440
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	227	ctatcaaggt	gacccagcac	ctgggaattg	ttggggagtg	caatgtgcag	tatgccttga	2040
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	233	ttgaggaggc	cttccagaag	gccctgcgca	tggtggatga	gaactgtgtg	ggctttgatc	2400
	234	acacagtgaa	accagtcage	gatatggagt	tggagactcc	aacagataag	cggatttttg	2460
	235	tggtggcagc	tgctttgtgg	gctggttatt	cagtggaccg	cctgtatgag	ctcacacgca	2520
	236	tcgaccgctg	gttcctgcac	cgaatgaagc	gtatcatcgc	acatgcccag	ctgctagaac	2580
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					gcacagagct			2700
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		_		_	ggggcaccac	_		2820
					tctaccgtat			2880
					tccgaaagat			2940
					actatgacat			3000
					tctatgagct			3060
				_	acatggccat			3.1.20
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					ccttcccctt			3720
					tcatggggga			3780
	<i>4</i> 3/	Laatgactgg	LECLYGAGTC	gegggagtaa	aggtgcctca	gitotocito	recedettaa	3840

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				tacctcaagg				3960
				attggcagct				4020
							gctgacttct	4080
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M>				cgtggcatcc				5580
				tccgacccag				5640
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				gcatctcccc				5760
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				cacctgttca				5880
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T.T				gacgtcgtcg				6180
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				gtggcctccc				6480
				actgatgtgc				6540
				gaagettget				6600
TAT				atggtggtga				6660
				ccccgngcag				6720
yy J				gccachgtgc gctgctgggc				6780
	200	yourullord	cccayyooda	gorgorggge	aayyaattee	agraceceer	acyggggcag	0/00

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309	tcttcattcc	tgcaccttaa	acctgtacag	tcatttttct	actgacttaa	taaacagccg	6960
310	agctgtccct	tg		1.5		•	6972

E--> 311 <210> SEQ ID NO: 3

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/658,659B

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Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 179

Skipped Sequences(NEW RULES):

Sequence(s)_missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

<400> sequence id number

Seq#:2

VERIFICATION SUMMARY

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Input Set : A:\11926-015001.txt

Output Set: N:\CRF4\11072002\I658659B.raw

```
L:46~M:258~W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
\text{L}:50~\text{M}:258~\text{W}: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
\text{L:}54~\text{M:}258~\text{W:} Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:58 M:258 W: Mandatory Feature missing < <220> Tag not found for SEQ ID#:1
L:62 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
L:63 M:341 W: (46) "n" or "Xaa" used, for SEQ:ID#:1 after pos.:240
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1080
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1200
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1320 L:87 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1680 L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:3120
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:3180
L:149 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:5400
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:5520
L:153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:5640
L:156 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:5820
L:157 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:5880
L:179 M:254 E: No. of Bases conflict, LENGTH: Input:2 Counted:7226 SEQ:1
L:179 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9
L:179 M:252 E: No. of Seq. differs <211> LENGTH:Input:7224 Found:7226 SEQ:1
L:193 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:193 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:193 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:1 differs:2
L:251 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:3420
L:265 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:4260
L:273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:4740
L:281 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5220
L:284 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5400 L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5460 L:290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:5760
L:296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6120
L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6600
L:305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:6660
L:311 M:214 E: (33) Seq.# missing, SEQ ID NO:2
L:325 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:328 M:341 W: (46) "n" or "Xaa" used; for SEQ ID#:3 after pos.:120
L:335 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:540 L:336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:600
L:354 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:1680 L:383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:3420
L:387 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:3660
L:388 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:3720
L:391 M:341 W: (46) "n" on "Xaa" used; for SEQ LD#:3 after pos::3900
L:407 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:411 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:415 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4 L:419 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:423 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/658,659B

DATE: 11/07/2002 TIME: 09:10:57

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L:426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:120
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:300
L:437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:780
L:441 M:341 W: (46) "n" or "Xaa" used for SEQ ID#:4 after pos.:1020
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:1320
L:457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:1980
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:2100
L:467 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:2580
L:468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:2640
L:486 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:490 M:258 W: Mandatory Feature missing <220> Tag not found for SEQ:ID#:5
L:498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:420
L:499 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:480
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:540
L:569 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:573 M:258 W: Mandatory Feature missing <220> Tag not found for SEQ ID#:6
L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:1020
L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:1080
L\!:\!598 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:1440
L:615 M:258 W: Mandatory Feature missing <220> Tag not found for SEQ ID#:7
L:619 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:623 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:240
L:629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:300
L:631 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:420
L:632 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:480
L:661 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:667 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:671 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:675 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:679 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:683 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:1009 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:1013 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:1017 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:1021 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:9
L:1079 M:258 W: Mandatory Feature missing <220> Tag not found for SEQ ID#:100
L:1083 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:1128 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1132 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1136 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1140 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1144 M:258 W: Mandatory Feature missing <220> Tag not found for SEQ ID#:11
L:1148 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1152 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:1266 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12 L:1318 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:13
\text{L:}1378~\text{M:}258~\text{W:} Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:1382 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
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L:1386 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:14
L:1445 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1449 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:1457 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:1461 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:1514 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:15
L:1518 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:1522 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:1526 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16
L:1526 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:16